## **AMENDMENTS TO THE CLAIMS**

All claim cancellations and amendments are done without prejudice.

Claims 1-214 (cancelled)

- 215. (Previously Presented) A method for producing oil with an altered fatty acid profile comprising extracting said oil with an altered fatty acid profile from a microbial cell culture produced by culturing a microbial cell comprising a recombinant nucleic acid that is a DNA molecule comprising the coding region of the sequence depicted in SEQ ID NO: 1, said nucleic acid operably linked to a promoter functional in said cell, wherein a polypeptide encoded by said nucleic acid is expressed in sufficient amount in said culture to alter the fatty acid profile.
- 216. (Previously presented) The method of claim 215, further comprising purifying a component of said oil.
- 217. (Previously presented) The method of claim 216, wherein said component is a phospholipid.
- 218. (Previously presented) The method of claim 216, wherein said component is a sulfolipid.
- 219. (Previously presented) The method of claim 216, wherein said component is a glycolipid.
- 220. (Previously presented) The method of claim 216, wherein said component is an acylglycerol.
- 221. (Previously presented) The method of claim 216, wherein said component is a monoacylglycerol.
- 222. (Previously presented) The method of claim 216, wherein said component is a diacylglycerol.
- 223. (Previously presented) The method of claim 216, wherein said component is a triacylglycerol.
- 224. (Previously presented) The method of claim 216, wherein said component is a fatty acid.

- 225. (Previously Presented) A method for producing oil with an altered fatty acid profile comprising extracting said oil with an altered fatty acid profile from a microbial cell culture produced by culturing a microbial cell comprising a recombinant nucleic acid that is a DNA molecule with at least 60% homology to the coding region of the sequence depicted in SEQ ID NO: 1, said nucleic acid operably linked to a promoter functional in said cell, wherein a polypeptide encoded by said nucleic acid forms a monounsaturated bond between carbons 6 and 7 of a fatty acid as numbered from a carboxy terminus thereof, wherein said polypeptide is expressed in sufficient amount in said culture to alter the fatty acid profile.
- 226. (Previously presented) The method of claim 225, further comprising purifying a component of said oil.
- 227. (Previously presented) The method of claim 226, wherein said component is a phospholipid.
- 228. (Previously presented) The method of claim 226, wherein said component is a sulfolipid.
- 229. (Previously presented) The method of claim 226, wherein said component is a glycolipid.
- 230. (Previously presented) The method of claim 226, wherein said component is an acylglycerol.
- 231. (Previously presented) The method of claim 226, wherein said component is a monoacylglycerol.
- 232. (Previously presented) The method of claim 226, wherein said component is a diacylglycerol.
- 233. (Previously presented) The method of claim 226, wherein said component is a triacylglycerol.
- 234. (Previously presented) The method of claim 226, wherein said component is a fatty acid.
- 235. (Currently Amended) A method for producing oil with an altered fatty acid profile comprising extracting said oil with an altered fatty acid profile from a microbial cell culture produced by culturing a microbial cell comprising a recombinant nucleic acid that is a DNA

molecule operably linked to a promoter functional in said cell to produce the microbial cell culture, wherein said nucleic acid <u>is encodes</u> a deletion mutant of the <u>nucleic acid-polypeptide</u> depicted in SEQ ID NO: <u>12</u>, wherein <u>a polypeptide encoded by said-nucleic acid-said deletion mutant</u> forms a monounsaturated bond between carbons 6 and 7 of a fatty acid as numbered from a carboxy terminus thereof, wherein said <u>polypeptide deletion mutant</u> is expressed in sufficient amount in said culture to alter the fatty acid profile.

- 236. (Previously presented) The method of claim 235, further comprising purifying a component of said oil.
- 237. (Previously presented) The method of claim 236, wherein said component is a phospholipid.
- 238. (Previously presented) The method of claim 236, wherein said component is a sulfolipid.
- 239. (Previously presented) The method of claim 236, wherein said component is a glycolipid.
- 240. (Previously presented) The method of claim 236, wherein said component is an acylglycerol.
- 241. (Previously presented) The method of claim 236, wherein said component is a monoacylglycerol.
- 242. (Previously presented) The method of claim 236, wherein said component is a diacylglycerol.
- 243. (Previously presented) The method of claim 236, wherein said component is a triacylglycerol.
- 244. (Previously presented) The method of claim 236, wherein said component is a fatty acid.

## Claims 245-254 (Cancelled)

255. (Previously presented) A method for producing oil with an altered fatty acid profile comprising extracting said oil with an altered fatty acid profile from a microbial cell culture produced by culturing a recombinant microbial cell comprising a polypeptide comprising the

- amino acid sequence depicted in SEQ ID NO:2, wherein said polypeptide is expressed in sufficient amount in said culture to alter the fatty acid profile.
- 256. (Previously presented) The method of claim 255, further comprising purifying a component of said oil.
- 257. (Previously presented) The method of claim 256, wherein said component is a phospholipid.
- 258. (Previously presented) The method of claim 256, wherein said component is a sulfolipid.
- 259. (Previously presented) The method of claim 256, wherein said component is a glycolipid.
- 260. (Previously presented) The method of claim 256, wherein said component is an acylglycerol.
- 261. (Previously presented) The method of claim 256, wherein said component is a monoacylglycerol.
- 262. (Previously presented) The method of claim 256, wherein said component is a diacylglycerol.
- 263. (Previously presented) The method of claim 256, wherein said component is a triacylglycerol.
- 264. (Previously presented) The method of claim 256, wherein said component is a fatty acid.
- 265. (Previously Presented) A method for producing oil with an altered fatty acid profile comprising extracting said oil with an altered fatty acid profile from a microbial cell culture produced by culturing a recombinant microbial cell comprising a polypeptide with at least 60% homology to the sequence depicted in SEQ ID NO: 2 to produce the microbial cell culture, wherein said polypeptide forms a monounsaturated bond between carbons 6 and 7 of a fatty acid as numbered from a carboxy terminus thereof, wherein said polypeptide is expressed in sufficient amount in said culture to alter the fatty acid profile.
- 266. (Previously presented) The method of claim 265, further comprising purifying a

component of said oil.

- 267. (Previously presented) The method of claim 266, wherein said component is a phospholipid.
- 268. (Previously presented) The method of claim 266, wherein said component is a sulfolipid.
- 269. (Previously presented) The method of claim 266, wherein said component is a glycolipid.
- 270. (Previously presented) The method of claim 266, wherein said component is an acylglycerol.
- 271. (Previously presented) The method of claim 266, wherein said component is a monoacylglycerol.
- 272. (Previously presented) The method of claim 266, wherein said component is a diacylglycerol.
- 273. (Previously presented) The method of claim 266, wherein said component is a triacylglycerol.
- 274. (Previously presented) The method of claim 266, wherein said component is a fatty acid.

Claims 275-297 (Cancelled)

298. (Previously Presented) A method for producing oil with an altered fatty acid profile comprising extracting said oil with an altered fatty acid profile from a microbial cell culture produced by culturing a microbial cell comprising a recombinant nucleic acid to produce the microbial cell culture, wherein said nucleic acid is a DNA molecule that hybridizes preferentially to a complement of the sequence depicted in SEQ ID NO: 1 under hybridization conditions suitable for selectively screening a recombinant DNA library using a probe comprising said complement, said recombinant DNA library comprising sequences obtained from a *Mortierella* species, said nucleic acid operably linked to a promoter functional in said cell, wherein a polypeptide encoded by said nucleic acid forms a monounsaturated bond between carbons 6 and 7 of a fatty acid as numbered from a carboxy terminus thereof, wherein said polypeptide is expressed in sufficient amount in said culture to alter the fatty acid profile.

- 299. (Previously Presented) The method of claim 298, wherein the *Mortierella* species is *Mortierella alpina*.
- 300. (Previously Presented) A method for producing oil with an altered fatty acid profile comprising extracting said oil with an altered fatty acid profile from a microbial cell culture produced by culturing a microbial cell, said microbial cell comprising a *Mortierella alpina*-derived means for forming a monounsaturated bond between carbons 6 and 7 of a fatty acid as numbered from a carboxy terminus thereof, wherein said means is used to produce said altered fatty acid profile.
- 301. (Previously Presented) The method of claim 299, further comprising purifying a component of said oil.
- 302. (Previously Presented) The method of claim 301, wherein said component is a phospholipid.
- 303. (Previously Presented) The method of claim 301, wherein said component is a sulfolipid.
- 304. (Previously Presented) The method of claim 301, wherein said component is a glycolipid.
- 305. (Previously Presented) The method of claim 301, wherein said component is an acylglycerol.
- 306. (Previously Presented) The method of claim 301, wherein said component is a monoacylglycerol.
- 307. (Previously Presented) The method of claim 301, wherein said component is a diacylglycerol.
- 308. (Previously Presented) The method of claim 301, wherein said component is a triacylglycerol.
- 309. (Previously Presented) The method of claim 301, wherein said component is a fatty acid.
- 310. (Previously Presented) The method of claim 300, further comprising purifying a component of said oil.

- 311. (Previously Presented) The method of claim 310, wherein said component is a phospholipid.
- 312. (Previously Presented) The method of claim 310, wherein said component is a sulfolipid.
- 313. (Previously Presented) The method of claim 310, wherein said component is a glycolipid.
- 314. (Previously Presented) The method of claim 310, wherein said component is an acylglycerol.
- 315. (Previously Presented) The method of claim 310, wherein said component is a monoacylglycerol.
- 316. (Previously Presented) The method of claim 310, wherein said component is a diacylglycerol.
- 317. (Previously Presented) The method of claim 310, wherein said component is a triacylglycerol.
- 318. (Previously Presented) The method of claim 310, wherein said component is a fatty acid.
- 319. (Previously Presented) The method of claim 225, wherein the recombinant nucleic acid has at least 80% homology to the sequence depicted in SEQ ID NO: 1.
- 320. (Previously Presented) The method of claim 265, wherein the polypeptide has at least 80% homology to the sequence depicted in SEQ ID NO: 2.
- 321. (Previously Presented) The method of claim 225, wherein the recombinant nucleic acid has at least 90% homology to the sequence depicted in SEQ ID NO: 1.
- 322. (Previously Presented) The method of claim 265, wherein the polypeptide has at least 90% homology to the sequence depicted in SEQ ID NO: 2.
- 323. (Previously Presented) The method of claim 225, wherein the recombinant nucleic acid has at least 95% homology to the sequence depicted in SEQ ID NO: 1.
- 324. (Previously Presented) The method of claim 265, wherein the polypeptide has at least 95% homology to the sequence depicted in SEQ ID NO: 2.

325. (Previously Presented) The method of claim 215, wherein said cell is a fungal cell. 326. (Previously Presented) The method of claim 225, wherein said cell is a fungal cell. 327. (Previously Presented) The method of claim 235, wherein said cell is a fungal cell. 328. (Previously Presented) The method of claim 255, wherein said cell is a fungal cell. 329. (Previously Presented) The method of claim 265, wherein said cell is a fungal cell. 330. (Previously Presented) The method of claim 299, wherein said cell is a fungal cell. 331. (Previously Presented) The method of claim 300, wherein said cell is a fungal cell. 332. (Previously Presented) The method of claim 319, wherein said cell is a fungal cell. 333. (Previously Presented) The method of claim 320, wherein said cell is a fungal cell. 334. (Previously Presented) The method of claim 321, wherein said cell is a fungal cell. 335. (Previously Presented) The method of claim 322, wherein said cell is a fungal cell. 336. (Previously Presented) The method of claim 323, wherein said cell is a fungal cell. 337. (Previously Presented) The method of claim 324, wherein said cell is a fungal cell. 338. (Previously Presented) The method of claim 325, wherein said fungal cell is a yeast cell. 339. (Previously Presented) The method of claim 326, wherein said fungal cell is a yeast cell. 340. (Previously Presented) The method of claim 327, wherein said fungal cell is a yeast cell. 341. (Previously Presented) The method of claim 328, wherein said fungal cell is a yeast cell. 342. (Previously Presented) The method of claim 329, wherein said fungal cell is a yeast cell. 343. (Previously Presented) The method of claim 330, wherein said fungal cell is a yeast cell. 344. (Previously Presented) The method of claim 331, wherein said fungal cell is a yeast

cell.

- 345. (Previously Presented) The method of claim 332, wherein said fungal cell is a yeast cell.
- 346. (Previously Presented) The method of claim 333, wherein said fungal cell is a yeast cell.
- 347. (Previously Presented) The method of claim 334, wherein said fungal cell is a yeast cell.
- 348. (Previously Presented) The method of claim 335, wherein said fungal cell is a yeast cell.
- 349. (Previously Presented) The method of claim 336, wherein said fungal cell is a yeast cell.
- 350. (Previously Presented) The method of claim 337, wherein said fungal cell is a yeast cell.
- 351. (Previously Presented) The method of claim 338, further comprising purifying a component of said oil.
- 352. (Previously Presented) The method of claim 339, further comprising purifying a component of said oil.
- 353. (Previously Presented) The method of claim 340, further comprising purifying a component of said oil.
- 354. (Previously Presented) The method of claim 341, further comprising purifying a component of said oil.
- 355. (Previously Presented) The method of claim 342, further comprising purifying a component of said oil.
- 356. (Previously Presented) The method of claim 343, further comprising purifying a component of said oil.
- 357. (Previously Presented) The method of claim 344, further comprising purifying a component of said oil.

- 358. (Previously Presented) The method of claim 345, further comprising purifying a component of said oil.
- 359. (Previously Presented) The method of claim 346, further comprising purifying a component of said oil.
- 360. (Previously Presented) The method of claim 347, further comprising purifying a component of said oil.
- 361. (Previously Presented) The method of claim 348, further comprising purifying a component of said oil.
- 362. (Previously Presented) The method of claim 349, further comprising purifying a component of said oil.
- 363. (Previously Presented) The method of claim 350, further comprising purifying a component of said oil.
- 364. (Currently Amended) The method of claim 351, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 365. (Currently Amended) The method of claim 352, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 366. (Currently Amended) The method of claim 353, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 367. (Currently Amended) The method of claim 354, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 368. (Currently Amended) The method of claim 355, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 369. (Currently Amended) The method of claim 356, wherein said component is selected

- from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 370. (Currently Amended) The method of claim 357, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 371. (Currently Amended) The method of claim 359, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 372. (Currently Amended) The method of claim 361, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.
- 373. (Currently Amended) The method of claim 363, wherein said component is selected from the group consisting of a phospholipids a phospholipid, a sulfolipid, a glycolipid, an acylglycerol, a monoacylglycerol, a diacylglycerol, a triacylglycerol, and a fatty acid.